



ENTERED

RAW SEQUENCE LISTING

4 <110> APPLICANT: Franklin, Richard L.

PATENT APPLICATION: US/09/938,269

DATE: 06/11/2002

TIME: 11:17:18

Input Set : A:\seqlist.txt

Output Set: N:\CRF3\06112002\1938269.raw

```
Cowling, Didier S.P.
         Hubbel, Jeffrey A.
         van de Wetering, Petra
  9 <120> TITLE OF INVENTION: Treatment of Trauma
 12 <130> FILE REFERENCE: 314572-105
 14 <140> CURRENT APPLICATION NUMBER: 09/938,269
 15 <141> CURRENT FILING DATE: 2001-08-23
17 <160> NUMBER OF SEQ ID NOS: 17
 19 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 300
 23 <212> TYPE: PRT
 24 <213> ORGANISM: Panaeu vanameii.
 26 <400> SEQUENCE: 1
 27 Leu Leu Ala Leu Val Ala Ala Ala Ser Ala Ala Glu Trp Arg Trp
                     5
                                        10
 29 Gln Phe Arg His Pro Thr Val Thr Pro Asn Pro Arg Ala Lys Asn Pro
 30
                20 -
31 Phe Arg Val Thr Lys Ser Ser Pro Val Gln Pro Pro Ala Val Arg Gly
       35
                              . 40
                                                 . 45
 33 Thr Lys Ala Val Glu Asn Cys Gly Pro Val Ala Pro Arg Asn Lys Ile
                           55
 35 Val Gly Gly Met Glu Val Thr Pro His Ala Tyr Pro Trp Gln Val Gly
                       70
 37 Leu Phe Ile Asp Asp Met Tyr Phe Cys Gly Gly Ser Ile Ile Ser Asp
                    85
                                        90
 39 Glu Trp Val Leu Thr Ala Ala His Cys Met Asp Gly Ala Gly Phe Val
               100
                                   105 ·
 41 Glu Val Val Met Gly Ala His Ser Ile His Asp Glu Thr Glu Ala Thr
                              . 120
                                                   125
 43 Gln Val Arg Ala Thr Ser Thr Asp Phe Phe Thr His Glu Asn Trp Asn
                           135
                                               140
45 Ser Phe Thr Leu Ser Asn Asp Leu Ala Leu Ile Lys Met Pro Ala Pro
                       150
                                           155
 47 Ile Glu Phe Asn Asp Val Ile Gln Pro Val Cys Leu Pro Thr Tyr Thr
                   165
                                       170
49 Asp Ala Ser Asp Asp Phe Val Gly Glu Ser Val Thr Leu Thr Gly Trp
               180
                                   185
 51 Gly Lys Pro Ser Asp Ser Ala Phe Gly Ile Ala Glu Gln Leu Arg Glu
     195
                           . 200
53 Val Asp Val Thr Thr Ile Thr Thr Ala Asp Cys Gln Ala Tyr Tyr Gly
54
       210
                           215
                                            . 220
```

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```
55 Ile Val Thr Asp Lys Ile Leu Cys Ile Asp Ser Glu Gly Gly His Gly
  56 225
                        230
                                           235
  57 Ser Cys Asn Gly Asp Ser Gly Gly Pro Met Asn Tyr Val Thr Gly Gly
                    245 .
                                    250
  59 Val Thr Gln Thr Arg Gly Ile Thr Ser Phe Gly Ser Ser Thr Gly Cys
               260
                                   265
  61 Glu Thr Gly Tyr Pro Asp Gly Tyr Thr Arg Val Thr Ser Tyr Leu Asp
  62 275 280
  63 Trp Ile Glu Ser Asn Thr Gly Ile Ala Ile Asp Pro
  64 290
                         295
  66 <210> SEQ ID NO: 2
  67 <211> LENGTH: 25
  68 <212> TYPE: PRT
  69 <213> ORGANISM: Panaeus vanameii
  71 <400> SEQUENCE: 2
  72 Ile Val Gly Gly Val Glu Ala Thr Pro His Ser Trp Pro His Gln Ala 🐭
               5
  73 1
                                       10
  74 Ala Leu Phe Ile Asp Asp Met Tyr Phe
       20
  77 <210> SEQ ID NO: 3
  78 <211> LENGTH: 20
  79 <212> TYPE: PRT
  80 <213> ORGANISM: Panaeus vanameii
  82 <220> FEATURE:
  83 <221> NAME/KEY: VARIANT
  84 <222> LOCATION: (1)...(20)
  85 <223> OTHER INFORMATION: Xaa = Any Amino Acid
  87 <400> SEQUENCE: 3
-> 88 Ile Val Gly Gly Val Glu Ala Thr Pro His Ser Xaa Pro His Gln Ala
  89 1
                                       10
  90 Ala Leu Phe Ile
  91
                20
  93 <210> SEQ ID NO: 4
  94 <211> LENGTH: 25
  95 <212> TYPE: PRT
  96 <213> ORGANISM: Panaeus monodon tryptic
  98 <400> SEQUENCE: 4
  99 Ile Val Gly Gly Thr Ala Val Thr Pro Gly Glu Phe Pro Tyr Gln Leu
          5 .
  100 1
                                        10
  101 Ser Phe Gln Asp Ser Ile Glu Gly Val
                 20
  104 <210> SEQ ID NO: 5
  105 <211> LENGTH: 25
  106 <212> TYPE: PRT
  107 <213> ORGANISM: Panaeus monodon chymotryptic
  109 <400> SEQUENCE: 5
  110 Ile Val Gly Val Glu Ala Val Pro Gly Val Trp Pro Tyr Gln Ala
  112 Ala Leu Phe Ile Ile Asp Met Tyr Phe
```

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                                                          TIME: 11:17:18
                    Input Set : A:\seqlist.txt
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     113
    115 <210> SEQ ID NO: 6
    116 <211> LENGTH: 25
    117 <212> TYPE: PRT
    118 <213> ORGANISM: Panaeus monodon chymotryptic
    120 <400> SEQUENCE: 6
    121 Ile Val Gly Gly Val Glu Ala Val Pro His Ser Trp Pro Tyr Gln Ala
                            10
    122 1
                        5
    123 Ala Leu Phe Ile Ile Asp Met Tyr Phe
    124
           . 20
    126 <210> SEQ ID NO: 7
    127 <211> LENGTH: 25
    128 <212> TYPE: PRT
    129 <213> ORGANISM: Uca pugilator enzyme
    131 <400> SEQUENCE: 7
    132 Ile Val Gly Gly Val Glu Ala Val Pro Asn Ser Trp Pro His Gln Ala
              5
    133 1
                                          10
    134 Ala Leu Phe Ile Asp Asp Met Tyr Phe
    135 20
    137 <210> SEQ ID NO: 8
    138 <211> LENGTH: 20
    139 <212> TYPE: PRT
    140 <213> ORGANISM: Uca pugilator enzyme
    142 <400> SEQUENCE: 8
    143 Ile Val Gly Gly Gln Asp Ala Thr Pro Gly Gln Phe Pro Tyr Gln Leu
                       .5 ..
                               10
                                                              15
    145 Ser Phe Gln Asp
                    20
    146
    148 <210> SEQ ID NO: 9
    149 <211> LENGTH: 20
    150 <212> TYPE: PRT
    151 <213> ORGANISM: Kamchatka crab
    153 <220> FEATURE:
    154 <221> NAME/KEY: VARIANT
    155 <222> LOCATION: (1)...(20)
    156 <223> OTHER INFORMATION: Xaa = Any Amino Acid
    158 <400> SEQUENCE: 9
W--> 159 Ile Val Gly Gly Gln Glu Ala Ser Pro Gly Ser Trp Pro Xaa Gln Val
    160 1
                                           10
    161 Gly Leu Phe Phe
    162
    164 <210> SEQ ID NO: 10
    165 <211> LENGTH: 20
    166 <212> TYPE: PRT
    167 <213> ORGANISM: Kamchatka crab
    169 <400> SEQUENCE: 10
    170 Ile Val Gly Gly Thr Glu Val Thr Pro Gly Glu Ile Pro Tyr Gln Leu
    171 1
    172 Ser Leu Gln Asp
```

RAW SEQUENCE LISTING

DATE: 06/11/2002

PATENT APPLICATION: US/09/938,269 TIME: 11:17:18 Input Set : A:\seqlist.txt Output Set: N:\CRF3\06112002\I938269.raw 173 175 <210> SEQ ID NO: 11 176 <211> LENGTH: 20 177 <212> TYPE: PRT 178 <213> ORGANISM: Kamchatka crab 180 <400> SEQUENCE: 11 181 Ile Val Gly Gly Thr Glu Val Thr Pro Gly Glu Ile Pro Tyr Gln Leu 182 5 10 183 Ser Phe Gln Asp 184 20 186 <210> SEQ ID NO: 12 187 <211> LENGTH: 20 188 <212> TYPE: PRT 189 <213> ORGANISM: Kamchatka crab 191 <220> FEATURE: 192 <221> NAME/KEY: VARIANT 193 <222> LOCATION: (1)...(20) 194 <223> OTHER INFORMATION: Xaa = Any Amino Acid 196 <400> SEQUENCE: 12 W--> 197 Ile Val Gly Gly Ser Glu Ala Thr Ser Gly Gln Phe Pro Tyr Gln Xaa 198 1 199 Ser Phe Gln Asp 200 20 202 <210> SEQ ID NO: 13 203 <211> LENGTH: 20 204 <212> TYPE: PRT 205 <213> ORGANISM: Crayfish protease 207 <400> SEQUENCE: 13 208 Ile Val Gly Gly Thr Asp Ala Thr Leu Gly Glu Phe Pro Tyr Gln Leu 209 5 1 210 Ser Phe Gln Asn 211 20 213 <210> SEQ ID NO: 14 214 <211> LENGTH: 25 215 <212> TYPE: PRT 216 <213> ORGANISM: Salmon enzyme 218 <400> SEQUENCE: 14 219 Ile Val Gly Gly Tyr Glu Cys Lys Ala Tyr Ser Gln Ala Tyr Gln Val 5 10 15 220 1 . 221 Ser Leu Asn Ser Gly Tyr His Tyr Cys 224 <210> SEQ ID NO: 15 225 <211> LENGTH: 25 226 <212> TYPE: PRT 227 <213> ORGANISM: Atlantic cod

230 Ile Val Gly Gly Tyr Glu Cys Thr Lys His Ser Gln Ala His Gln Val

RAW SEQUENCE LISTING

229 <400> SEQUENCE: 15

232 Ser Leu Asn Ser Gly Tyr His Tyr Cys

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Input Set : A:\seqlist.txt
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233	20		25
235	<210> SEQ ID NO: 16		
236	<211> LENGTH: 25	4	
237	<212> TYPE: PRT		00
238	<213> ORGANISM: Atla	antic cod	
240	<400> SEQUENCE: 16	* *	
241	Ile Val Gly Gly Tyr	Glu Cys Thr	Arg His Ser Gln Ala His Gln Val
242	. 5		10 15
243.	Ser Leu Asn Ser Gly	Tyr His Tyr	Cys
244	20		25
246	<210> SEQ ID NO: 17		
247	<211> LENGTH: 37		
248	<212> TYPE: PRT		
249	<213> ORGANISM: Atla	intic cod	
	<400> SEQUENCE: 17	•	
252	Ile Val Gly Gly Tyr	Gln Cys Glu	ı Ala His Ser Gln Ala His Gln Val
253	1 5	• 1	10 15
254	Ser Leu Asn Ser Gly	Tyr His Tyr	Cys Gly Gly Ser Leu Ile Asn Trp
255	20		25 30
-	Val Val Ser Ala Ala		
257	35		

RAW SEQUENCE LISTING ERROR SUMMARY

DATE: 06/11/2002 TIME: 11:17:19

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Input Set : A:\seqlist.txt

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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; Xaa Pos. 12 Seq#:9; Xaa Pos. 14 Seq#:12; Xaa Pos. 16 VERIFICATION SUMMARY

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Input Set : A:\seqlist.txt

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L:88 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0 L:159 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0 L:197 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0